

SEQUENCE LISTING

<110> Ghayur, Tariq et al.

<120> ANTIBODIES THAT BIND HUMAN INTERLEUKIN-18 AND METHODS
OF MAKING AND USING

<130> BBI-149

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<150> 60/181,608

<151> 2000-02-10

<160> 71

<170> PatentIn Ver. 2.1

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Met Thr Asp Ser Asp Cys Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile
35 40 45

Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
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Ser Val Lys Cys Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile
65 70 75 80

Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
85 90 95

Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys
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Met Gln Phe Glu Ser Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
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35 40 45

Gly Glu Glu Ser Asn Asp Lys Ile Pro Val Ala Leu Gly Leu Lys Glu
50 55 60

Lys Asn Leu Tyr Leu Ser Cys Val Leu Lys Asp Asp Lys Pro Thr Leu
65 70 75 80

Gln Leu Glu Ser Val Asp Pro Lys Asn Tyr Pro Lys Lys Lys Met Glu
85 90 95

Lys Arg Phe Val Phe Asn Lys Ile Glu Ile Asn Asn Lys Leu Glu Phe
100 105 110

Glu Ser Ala Gln Phe Pro Asn Trp Tyr Ile Ser Thr Ser Gln Ala Glu
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Ala Leu Phe Leu Gly Ile His Gly Gly Lys Met Cys Leu Ser Cys Val
50 55 60
Lys Ser Gly Asp Glu Thr Arg Leu Gln Leu Glu Ala Val Asn Ile Thr
65 70 75 80
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Ser Asp Ser Gly Pro Thr Thr Ser Phe Glu Ser Ala Ala Cys Pro Gly
100 105 110
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35 40 45
Pro Arg Ser Ser Ser Arg Ile Ala Leu His Asp Cys Val Leu Glu Phe
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Trp Pro Val Glu Leu Asn Asp Thr Gly Ser Tyr Phe Phe Gln Met Lys
65 70 75 80
Asn Tyr Thr Gln Lys Trp Lys Leu Asn Val Ile Arg Arg Asn Lys His

85										90										95									
Ser	Cys	Phe	Thr	Glu	Arg	Gln	Val	Thr	Ser	Lys	Ile	Val	Glu	Val	Lys														
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Lys	Phe	Phe	Gln	Ile	Thr	Cys	Glu	Asn	Ser	Tyr	Tyr	Gln	Thr	Leu	Val														
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Tyr	Tyr	Ser	Cys	Val	His	Phe	Leu	His	His	Asn	Gly	Lys	Leu	Phe	Asn														
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Ile	Thr	Lys	Thr	Phe	Asn	Ile	Thr	Ile	Val	Glu	Asp	Arg	Ser	Asn	Ile														
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Val	Pro	Val	Leu	Leu	Gly	Pro	Lys	Leu	Asn	His	Val	Ala	Val	Glu	Leu														
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Gly	Lys	Asn	Val	Arg	Leu	Asn	Cys	Ser	Ala	Leu	Leu	Asn	Glu	Glu	Asp														
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Val	Ile	Tyr	Trp	Met	Phe	Gly	Glu	Glu	Asn	Gly	Ser	Asp	Pro	Asn	Ile														
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His	Glu	Glu	Lys	Glu	Met	Arg	Ile	Met	Thr	Pro	Glu	Gly	Lys	Trp	His														
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Val	Leu	Tyr	Asn	Cys	Thr	Val	Ala	Ser	Thr	Gly	Gly	Thr	Asp	Thr	Lys														
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 35 40 45
 Ala Ser Arg Ile His Gln His Lys Glu Lys Leu Trp Phe Val Pro Ala
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 Lys Val Glu Asp Ser Gly His Tyr Tyr Cys Val Val Arg Asn Ser Ser

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Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn
50 55 60

Ser Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp
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35 40 45

Ile Tyr Gly Asn Asp Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser
50 55 60

Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln
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35 40 45

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Tyr Tyr Ala Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn
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Ser Lys Asn Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp
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Thr Ala Val Tyr Tyr Cys Ala Arg Asp Asp Asp Asp Tyr Asp Phe Asp
          85          90          95
Tyr Trp Gly Arg Gly Thr Met Val Thr Val Ser Ser Gly Gly Gly Gly
  100          105          110
Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Ala Gln Ser Val Leu
  115          120          125
Thr Gln Pro Pro Ser Ala Ser Gly Thr Pro Gly Gln Arg Val Thr Ile
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Ser Cys Ser Gly Ser Ser Ser Asn Ile Gly Ile Asn Ala Val Asn Trp
  145          150          155          160
Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu Ile Tyr Gly Asn
          165          170          175
Asp Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Lys Ser
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Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln Ser Gly Asp Glu
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Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly
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Asn Ala Pro Arg Thr Ile Phe Ile Ile Ser Met Tyr Lys Asp
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Met Thr Asp Ser Asp Cys Arg Asp Asn Ala Pro Arg Thr Ile Phe Ile
 35 40 45

Ile Ser Met Tyr Lys Asp Ser Gln Pro Arg Gly Met Ala Val Thr Ile
 50 55 60

Ser Val Lys Cys Glu Lys Ile Ser Thr Leu Ser Cys Glu Asn Lys Ile
 65 70 75 80

Ile Ser Phe Lys Glu Met Asn Pro Pro Asp Asn Ile Lys Asp Thr Lys
 85 90 95

Ser Asp Ile Ile Phe Phe Gln Arg Ser Val Pro Gly His Asp Asn Lys
 100 105 110

Met Gln Phe Glu Ser Ser Ser Tyr Glu Gly Tyr Phe Leu Ala Cys Glu
 115 120 125

Lys Glu Arg Asp Leu Phe Lys Leu Ile Leu Lys Lys Glu Asp Glu Leu
130 135 140

Gly Asp Arg Ser Ile Met Phe Thr Val Gln Asn Glu Asp
145 150 155

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Ser Met Lys Val Ser Cys Lys Thr Ser Gly Tyr Thr Phe Thr Gly Tyr
20 25 30

tat atc cac tgg gtg cga cag gcc cct gga cag gga ttc gag tgg ata 144
Tyr Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Phe Glu Trp Ile
35 40 45

gga cgg ctc aac ccc acc act ggt gac gca aat ttt gca gaa aag ttt 192
Gly Arg Leu Asn Pro Thr Thr Gly Asp Ala Asn Phe Ala Glu Lys Phe
50 55 60

cag ggc agg gtc gcc ctg acc aga gac acg tcc atc agc aca gcc tat 240
Gln Gly Arg Val Ala Leu Thr Arg Asp Thr Ser Ile Ser Thr Ala Tyr
65 70 75 80

tta caa cta gac agc ctc aaa tct gac gac acg gcc gta tat tat tgt 288
Leu Gln Leu Asp Ser Leu Lys Ser Asp Asp Thr Ala Val Tyr Tyr Cys
85 90 95

gcg gga aaa gag ggt gcc tgg gcc cag ggc acc ctg gtc acc gtc tcg 336
Ala Gly Lys Glu Gly Ala Trp Gly Gln Gly Thr Leu Val Thr Val Ser
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Ser

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20 25 30

Tyr Ile His Trp Val Arg Gln Ala Pro Gly Gln Gly Phe Glu Trp Ile
 35 40 45
 Gly Arg Leu Asn Pro Thr Thr Gly Asp Ala Asn Phe Ala Glu Lys Phe
 50 55 60
 Gln Gly Arg Val Ala Leu Thr Arg Asp Thr Ser Ile Ser Thr Ala Tyr
 65 70 75 80
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 Ala Gly Lys Glu Gly Ala Trp Gly Gln Gly Thr Leu Val Thr Val Ser
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 Ser

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 Thr Val Arg Ile Thr Cys Gln Gly Asp Ser Leu Arg His Phe Tyr Pro
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 aac tgg tac cag cag aag cca gga cag gcc cct gta ctt gtc atc tat 144
 Asn Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
 35 40 45
 ggt aaa aac aat cgg ccc tca ggg atc cca gac cga ttc tct ggc tcc 192
 Gly Lys Asn Asn Arg Pro Ser Gly Ile Pro Asp Arg Phe Ser Gly Ser
 50 55 60
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 Gly Ser Gly Asn Thr Gly Ser Leu Thr Ile Thr Gly Ala Gln Ala Glu
 65 70 75 80
 gat gag gct gac tat tac tgt ggc tcc cgg gac agc agt ggt atc cat 288
 Asp Glu Ala Asp Tyr Thr Cys Gly Ser Arg Asp Ser Ser Gly Ile His
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 gtg gta ttc ggc gga ggg acc aag gtc acc gtc cta ggt 327
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Ala Arg Asp Asp Asp Asp Tyr Asp Phe Asp Tyr Trp Gly Arg Gly Thr

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Met Val Thr Val Ser Ser			
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Ala Met Ser Ser Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu Trp Val			
35	40	45	
Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp Ser Val			
50	55	60	
Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu Tyr			
65	70	75	80
Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr Cys			
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Arg Val Thr Ile Ser Cys Ser Gly Ser Ser Ser Asn Ile Gly Ile Asn			
20	25	30	
gct gta aac tgg tac cag cag ctc cca gga acg gcc ccc aaa ctc ctc			144
Ala Val Asn Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu			
35	40	45	
atc tat ggt aat gat cag ccg ccc tca ggg gtc cct gac cga ttc tct			192
Ile Tyr Gly Asn Asp Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser			

50	55	60	
ggc tcc aag tct ggc acc tca gcc tcc ctg gcc atc agt ggg ctc cag			240
Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln			
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tct gag gat gag gct gat tat aac tgt gca gca tgg gat gac agc ctg			288
Ser Glu Asp Glu Ala Asp Tyr Asn Cys Ala Ala Trp Asp Asp Ser Leu			
85	90	95	
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Ser Gly Pro Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly			
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20 25 30
Ala Val Asn Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
35 40 45
Ile Tyr Gly Asn Asp Gln Arg Pro Ser Gly Val Pro Asp Arg Phe Ser
50 55 60
Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Leu Gln
65 70 75 80
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85 90 95
Ser Gly Pro Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly
100 105 110

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